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Robust radiogenomics approach to the identification of *EGFR* mutations among patients with NSCLC from three different countries using topologically invariant Betti numbers

Kenta Ninomiya¹, Hidetaka Arimura²*, Wai Yee Chan³*, Kentaro Tanaka⁴, Shinichi Mizuno⁵, Nadia Fareeda Muhammad Gowdh³, Nur Adura Yaakup³, Chong-Kin Liam⁶, Chee-Shee Chai⁷, Kwan Hoong Ng³

1 Division of Medical Quantum Science, Department of Health Sciences, Graduate School of Medical Sciences, Kyushu University, Fukuoka, Japan, 2 Faculty of Medical Sciences, Division of Medical Quantum Science, Department of Health Sciences, Kyushu University, Fukuoka, Japan, 3 Faculty of Medicine, Department of Biomedical Imaging, University of Malaya, Kuala Lumpur, Malaysia, 4 Department of Respiratory Medicine, Kyushu University Hospital, Fukuoka, Japan, 5 Division of Medical Sciences and Technology, Department of Health Sciences, Graduate School of Medical Sciences, Kyushu University, Fukuoka, Japan, 6 Faculty of Medicine, Department of Medicine, University of Malaya, Kuala Lumpur, Malaysia, 7 Faculty of Medicine and Health Science, Department of Medicine, University Malaysia Sarawak, Kota Samarahan, Sarawak, Malaysia

* arimurah@med.kyushu-u.ac.jp (HA); waiyeec@ummc.edu.my (WYC)

Abstract

Objectives

To propose a novel robust radiogenomics approach to the identification of epidermal growth factor receptor (*EGFR*) mutations among patients with non-small cell lung cancer (NSCLC) using Betti numbers (BNs).

Materials and methods

Contrast enhanced computed tomography (CT) images of 194 multi-racial NSCLC patients (79 *EGFR* mutants and 115 wildtypes) were collected from three different countries using 5 manufacturers' scanners with a variety of scanning parameters. Ninety-nine cases obtained from the University of Malaya Medical Centre (UMMC) in Malaysia were used for training and validation procedures. Forty-one cases collected from the Kyushu University Hospital (KUH) in Japan and fifty-four cases obtained from The Cancer Imaging Archive (TCIA) in America were used for a test procedure. Radiomic features were obtained from BN maps, which represent topologically invariant heterogeneous characteristics of lung cancer on CT images, by applying histogram- and texture-based feature computations. A BN-based signature was determined using support vector machine (SVM) models with the best combination of features that maximized a robustness index (RI) which defined a higher total area under receiver operating characteristics curves (AUCs) and lower difference of AUCs